

Package ‘fonctionr’

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Type Package

Title Easy Estimation and Vizualisation of Indicators from Data with Complex Design

Version 0.5.1

Description Many functions to easily vizualise and estimate indicators such as proportions, means, medians and continuous/discrete distributions from complex survey data. The package also estimates confidence intervals for all indicators, compares different groups and computes different statistical tests.

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URL <https://jgires.github.io/fonctionr/>,
<https://github.com/jgires/fonctionr/>

BugReports <https://github.com/jgires/fonctionr/issues>

Depends R (>= 4.1.0)

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central_group	<i>central_group</i>
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Description

Function to compare means or medians among different groups based on complex survey data. It produces a list containing a table, including the confidence intervals of the indicators, a ready-to-be published ggplot graphic and a statistical test.

In case of mean comparison, the statistical test is a Wald test (using `survey::regTermTest()`). In case of median comparison the statistical test is a Kruskal Wallis test (using `survey::svyranktest()`). The confidence intervals and the statistical test are taking into account the complex survey design. In case of facets, the statistical test is computed on the total means or medians between facets (and not within facets). In case of second group (`group.fill`), no statistical test is computed.

Exporting the results to an Excell file is possible.

Usage

```
central_group(
  data,
  group,
  quanti_exp,
  type,
  group.fill = NULL,
  facet = NULL,
  filter_exp = NULL,
  ...,
)
```

```
na.rm.group = TRUE,  
na.rm.facet = TRUE,  
total = TRUE,  
reorder = FALSE,  
show_ci = TRUE,  
show_n = FALSE,  
show_value = TRUE,  
show_labs = TRUE,  
total_name = NULL,  
digits = 0,  
unit = "",  
dec = NULL,  
col = NULL,  
pal = "OBSS_Spring",  
direction = 1,  
desaturate = 0,  
lighten = 0,  
darken = 0,  
dodge = 0.9,  
font = "Roboto",  
wrap_width_y = 25,  
wrap_width_leg = 25,  
legend_ncol = 4,  
title = NULL,  
subtitle = NULL,  
xlab = NULL,  
ylab = NULL,  
legend_lab = NULL,  
caption = NULL,  
lang = "fr",  
theme = "fonctionr",  
coef_font = 1,  
export_path = NULL  
)  
  
median_group(..., type = "median")  
  
mean_group(..., type = "mean")
```

Arguments

data	A dataframe or an object from the survey package or an object from the srvyr package.
group	A variable defining groups to be compared.
quanti_exp	An expression defining the quantitative variable from which the mean/median is computed. Notice that if any observations with NA in at least one of the variable in quanti_exp are excluded for the computation of the indicators.
type	"mean" to compute mean by group ; "median" to compute median by group.

<code>group.fill</code>	A variable defining a second variable of groups to be compared.
<code>facet</code>	A variable defining the faceting group.
<code>filter_exp</code>	An expression filtering the data, preserving the design. Notice that <code>filter_exp</code> works as <code>srvyr::filter()</code> : it excludes observations for which <code>filter_exp</code> results into NA. It is often the case when NA is present on one of the filter variables.
<code>...</code>	All options possible in <code>srvyr::as_survey_design()</code> .
<code>na.rm.group</code>	TRUE if you want to remove observations with NA on the group and the <code>group.fill</code> variables. FALSE if you want to create a group with the NA values for the group variable and a <code>group.fill</code> with the NA values for the <code>group.fill</code> variable. Default is TRUE.
<code>na.rm.facet</code>	TRUE if you want to remove observations with NA on the facet variable. FALSE if you want to create a facet with the NA values for the facet variable. Default is TRUE.
<code>total</code>	TRUE if you want to compute a total, FALSE if you don't. The default is TRUE.
<code>reorder</code>	TRUE if you want to reorder the groups according to the mean/median. NA value, if <code>na.rm.group = FALSE</code> , is not included in the reorder. In case of facets, the groups are reordered based on each median group. Default is FALSE.
<code>show_ci</code>	TRUE if you want to show the error bars on the graphic. FALSE if you don't want to show the error bars. Default is TRUE.
<code>show_n</code>	TRUE if you want to show on the graphic the number of observations in the sample in each group. FALSE if you don't want to show this number. Default is FALSE.
<code>show_value</code>	TRUE if you want to show the mean/median in each group on the graphic. FALSE if you don't want to show the mean/median. Default is TRUE.
<code>show_labs</code>	TRUE if you want to show axes and legend (in case of a <code>group.fill</code>) labels. FALSE if you don't want to show any labels on axes and legend. Default is TRUE.
<code>total_name</code>	Name of the total displayed on the graphic. Default is "Total" in French and in English and "Totaal" in Dutch.
<code>digits</code>	Number of decimal places displayed on the values labels on the graphic. Default is 0.
<code>unit</code>	Unit displayed on the graphic. Default is none ("").
<code>dec</code>	Decimal mark displayed on the graphic. Default depends on lang: ", " for fr and nl ; "." for en.
<code>col</code>	Color of the bars if there is no <code>group.fill</code> . <code>col</code> must be a R color or an hexadecimal color code. Default color used depends on type : "deeeppink3" for mean and "mediumorchid3" for median. The colors of total and NA group (in case of <code>na.rm.group = FALSE</code>) are always "grey40" and "grey". If there is a <code>group.fill</code> , <code>col</code> has no effect and <code>pal</code> argument should be used instead.
<code>pal</code>	Colors of the bars if there is a <code>group.fill</code> . <code>pal</code> must be vector of R colors or hexadecimal colors or a palette from packages MetBrewer or PrettyCols or a palette from fonctionr. The color of missing values for <code>group.fill</code> (in case of <code>na.rm.group = FALSE</code>) and for the total are always "grey" and "grey40". If there is no <code>group.fill</code> , <code>pal</code> has no effect and <code>col</code> argument should be used instead.

direction	Direction of the palette color. Default is 1. The opposite direction is -1. If there is no <code>group.fill</code> , this argument has no effect.
desaturate	Numeric specifying the amount of desaturation where 1 corresponds to complete desaturation (no colors, grey layers only), 0 to no desaturation, and values in between to partial desaturation. Default is 0. It affects only the palette (<code>pal</code> , if there is a second group) and not the monocolour (<code>col</code> , if there is no second group). See <code>colorspace::desaturate</code> function from <code>colorspace</code> package for details. If <code>desaturate</code> and <code>lighten/darken</code> arguments are used, <code>lighten/darken</code> is applied in a second time (i.e. on the color transformed by <code>desaturate</code>).
lighten	Numeric specifying the amount of lightening. Negative numbers cause darkening. Value should be ranged between -1 (black) and 1 (white). Default is 0. It doesn't affect the color of NA (in case of <code>na.rm.group = FALSE</code>). It affects only the palette (<code>pal</code> , if there is a second group) and not the monocolour (<code>col</code> , if there is no second group). See <code>colorspace::desaturate</code> for details. If both argument <code>lighten</code> and <code>darken</code> are used (not advised), <code>darken</code> is applied in a second time (i.e. on the color transformed by <code>lighten</code>).
darken	Numeric specifying the amount of lightening. Negative numbers cause lightening. Value should be ranged between -1 (white) and 1 (black). Default is 0. It doesn't affect the color of NA (in case of <code>na.rm.group = FALSE</code>). It affects only the palette (<code>pal</code> , if there is a second group) and not the monocolour (<code>col</code> , if there is no second group). See <code>colorspace::desaturate</code> for details. If both argument <code>lighten</code> and <code>darken</code> are used (not advised), <code>darken</code> is applied in a second time (i.e. on the color transformed by <code>lighten</code>).
dodge	Width of the bars. Default is 0.9 to let a small space between bars. A value of 1 leads to no space between bars. Values higher than 1 are not advised because they cause an overlapping of the bars. <code>dodge</code> doesn't affect the spaces between second groups (<code>group.fill</code>). There is always no space between second groups.
font	Font used in the graphic. See <code>load_and_active_fonts()</code> for available fonts. Default is "Roboto".
wrap_width_y	Number of characters before going to the line for the labels of the groups. Default is 25.
wrap_width_leg	Number of characters before going to the line for the labels of the <code>group.fill</code> . Default is 25.
legend_ncol	Number of columns in the legend. Default is 4.
title	Title of the graphic.
subtitle	Subtitle of the graphic.
xlab	X label on the graphic. As <code>ggplot2::coord_flip()</code> is used in the graphic, <code>xlab</code> refers to the x label on the graphic, after the <code>ggplot2::coord_flip()</code> , and not to the x variable in the data. Default (<code>xlab = NULL</code>) displays, for <code>type = "mean"</code> , "Moyenne : " (if <code>lang = "fr"</code>), "Mean:" (if <code>lang = "en"</code>) or "Gemiddelde:" (if <code>lang = "nl"</code>), or, for <code>type = "median"</code> , "Médiane : " (if <code>lang = "fr"</code>), "Median:" (if <code>lang = "en"</code>) or "Mediaan:" (if <code>lang = "nl"</code>), followed by the <code>quanti_exp</code> argument. To show no X label, use <code>xlab = ""</code> .
ylab	Y label on the graphic. As <code>ggplot2::coord_flip()</code> is used in the graphic, <code>ylab</code> refers to the y label on the graphic, after the <code>ggplot2::coord_flip()</code> ,

	and not to the y variable in the data. Default (<code>ylab = NULL</code>) displays the name of the group variable. To show no Y label, use <code>ylab = ""</code> .
<code>legend_lab</code>	Legend (fill) label on the graphic. If <code>legend_lab = NULL</code> , legend label on the graphic will be <code>group.fill</code> . To show no legend label, use <code>legend_lab = ""</code> .
<code>caption</code>	Caption of the graphic. This caption goes under the default caption showing the result of the statistical test. There is no way of not showing the result of the chi-square test as a caption.
<code>lang</code>	Language of the indications on the graphic. Possibilities are "fr" (french), "nl" (dutch) and "en" (english). Default is "fr".
<code>theme</code>	Theme of the graphic. Default is "fonctionr". "IWEPS" adds y axis lines and ticks. NULL uses the default grey ggplot2 theme.
<code>coef_font</code>	A multiplier factor for font size of all fonts on the graphic. Default is 1. Useful when exporting the graphic for a publication (e.g. in a Quarto document).
<code>export_path</code>	Path to export the results in an xlsx file. The file includes three (without <code>group.fill</code>) or two sheets (with a <code>group.fill</code>): the table, the graphic and the statistical test result.

Value

A list that contains a table, a ggplot graphic and, in most cases, a statistical test.

Examples

```
# Loading of data
data(eusilc, package = "laeken")

# Creation of age categories
eusilc$age_cat <- cut(eusilc$age,
breaks = 6,
include.lowest = TRUE)

# Calculation of income means by age category with fonctionr, taking sample design into account
eusilc_mean <- mean_group(
  eusilc,
  group = age_cat,
  quanti_exp = eqIncome / 12,
  strata = db040,
  ids = db030,
  weight = rb050,
  title = "Mean of equivalised income in household by age of individuals",
  subtitle = "Example with austrian SILC data from 'laeken' package"
)

# Results in graph form
eusilc_mean$graph

# Results in table format
eusilc_mean$tab
```

distrib_continuous	<i>distrib_continuous</i>
--------------------	---------------------------

Description

Function to describe the distribution of a continuous variable from complex survey data. It produces a list containing a density table (`dens`), a central value table (`tab`), a quantile table (`quant`) and a ready-to-be published ggplot graphic (`graph`).

The density table contains x-y coordinates to draw a density curve. The central value table contains the median or the mean of the continuous variable, with its confidence interval, the sample size and the estimation of the total, with its confidence interval. The quantile table contains quantiles and their confidence intervals. The quantiles and the limits are used as thicks on the X axe of the graphic. The confidence intervals are taking into account the complex survey design.

Exporting those results to an Excell file is possible.

Usage

```
distrib_continuous(  
  data,  
  quanti_exp,  
  type = "median",  
  facet = NULL,  
  filter_exp = NULL,  
  ...,  
  na.rm.facet = TRUE,  
  quantiles = seq(0.1, 0.9, 0.1),  
  bw = 1,  
  resolution = 1024,  
  limits = NULL,  
  show_mid_line = TRUE,  
  show_ci_lines = TRUE,  
  show_ci_area = FALSE,  
  show_quant_lines = FALSE,  
  show_n = FALSE,  
  show_value = TRUE,  
  show_labs = TRUE,  
  digits = 0,  
  unit = "",  
  dec = NULL,  
  pal = NULL,  
  col_density = c("#00708C", "mediumturquoise"),  
  color = NULL,  
  col_border = NA,  
  font = "Roboto",  
  title = NULL,  
  subtitle = NULL,
```

```

xlab = NULL,
ylab = NULL,
caption = NULL,
lang = "fr",
theme = "fonctionr",
coef_font = 1,
export_path = NULL
)

distrib_c(...)

```

Arguments

<code>data</code>	A dataframe or an object from the survey package or an object from the <code>srvyr</code> package.
<code>quanti_exp</code>	An expression defining the quantitative variable the variable to be described. Notice that any observations with NA in at least one of the variable in <code>quanti_exp</code> are excluded for the computation of the density and of the indicators.
<code>type</code>	Type of central value : "mean" to compute mean as the central value ; "median" to compute median as the central value.
<code>facet</code>	Not yet implemented.
<code>filter_exp</code>	An expression filtering the data, preserving the design. Notice that <code>filter_exp</code> works as <code>srvyr::filter()</code> : it excludes observations for which <code>filter_exp</code> results into NA. It is often the case when NA is present on one of the filter variables.
<code>...</code>	All options possible in <code>srvyr::as_survey_design()</code> .
<code>na.rm.facet</code>	Not yet implemented.
<code>quantiles</code>	Quantiles computed. Default are deciles.
<code>bw</code>	The smoothing bandwidth to be used. The kernels are scaled such that this is the standard deviation of the smoothing kernel. Default is 1.
<code>resolution</code>	Resolution of the density curve. Default is 1024.
<code>limits</code>	Limits of the X axe of the graphic. Does not apply to the computation of indicators (median/mean and quantiles). Default is NULL to show the entire distribution on the graphic.
<code>show_mid_line</code>	TRUE if you want to show the mean or median (depending on type) as a line on the graphic. FALSE if you do not want to show it. Default is TRUE.
<code>show_ci_lines</code>	TRUE if you want to show confidence interval of the mean or median (depending on type) as dotted lines on the graphic. FALSE if you do not want to show it as lines. Default is TRUE.
<code>show_ci_area</code>	TRUE if you want to show confidence interval of the mean or median (depending on type) as a coloured area on the graphic. FALSE if you do not want to show it as an area. Default is FALSE.
<code>show_quant_lines</code>	TRUE if you want to show quantiles as lines on the graphic. FALSE if you do not want to show them as lines. Default is FALSE.

show_n	TRUE if you want to show on the graphic the number of individuals in the sample in each quantile. FALSE if you do not want to show the numbers. Default is FALSE.
show_value	TRUE if you want to show the value of the mean/median (depending on type) on the graphic. FALSE if you do not want to show the mean/median. Default is TRUE.
show_labs	TRUE if you want to show axes labels. FALSE if you do not want to show any labels on axes. Default is TRUE.
digits	Number of decimal places displayed on the values labels on the graphic. Default is 0.
unit	Unit displayed on the graphic. Default is none ("").
dec	Decimal mark shown on the graphic. Depends on lang: ", " for fr and nl ; "." for en.
pal	For compatibility with older versions.
col_density	Color of the density area. It may be one color or a vector with several colors. Colors should be R color or an hexadecimal color code. In case of one color, the density is monocolour. In case of a vector, the quantile areas are painted in continuous colors going from the last color in the vector (center quantile) to the first color (first and last quantiles). In case of an even quantile area numbers (e.g. deciles, quartiles) the last color of the vector is only applied to the highcenter quantile area to avoid two continuous quantile areas having the same color.
color	Not currently used except for compatibility with old versions.
col_border	Color of the density line. Color should be one R color or one hexadecimal color code. Default (NULL) does not draw the density line.
font	Font used in the graphic. See load_and_active_fonts() for available fonts. Default is "Roboto".
title	Title of the graphic.
subtitle	Subtitle of the graphic.
xlab	X label on the graphic. If xlab = NULL, X label on the graphic will be quanti_exp.
ylab	Y label on the graphic. If ylab = NULL, Y label on the graphic will be "Densité" (if lang = "fr"), "Density" (if lang = "en") or "Densiteit" (if lang = "nl").
caption	Caption of the graphic.
lang	Language of the indications on the graphic. Possibilities are "fr" (french), "nl" (dutch) and "en" (english). Default is "fr".
theme	Theme of the graphic. Default is "fonctionr". "IWEPS" adds y axis lines and ticks. NULL uses the default grey ggplot2 theme.
coef_font	A multiplier factor for font size of all fonts on the graphic. Default is 1. Usefull when exporting the graphic for a publication (e.g. in a Quarto document).
export_path	Path to export the results in an xlsx file. The file includes four sheets: the central value table, the quantile table, the density table and the graphic.

Value

A list that contains a density table (dens), a central value table (tab), a quantile table (quant) and a ggplot graphic (graph).

Examples

```

# Loading of data
data(eusilc, package = "laeken")

# Computation, taking sample design into account
eusilc_dist_c <- distrib_c(
  eusilc,
  quanti_exp = eqIncome,
  strata = db040,
  ids = db030,
  weight = rb050,
  limits = c(0, 50000),
  title = "Distribution of eq. income",
  subtitle = "Example with austrian SILC data from 'laeken' package"
)

# Results in graph form
eusilc_dist_c$graph

# Results in table format
eusilc_dist_c$tab

```

distrib_discrete *distrib_discrete*

Description

Function to describe the distribution of a discrete variable from complex survey data.

It produces a list containing a table, including the confidence intervals of the indicators, a ready-to-be published ggplot graphic and, if proportions for H0 are specified, a Chi-Square statistical test (using `survey::svyqofchisq()`). The confidence intervals and the statistical test are taking into account the complex survey design. In case of facets, no statistical test is (yet) computed.

Exporting those results to an Excell file is possible.

Usage

```

distrib_discrete(
  data,
  quali_var,
  facet = NULL,
  filter_exp = NULL,
  ...,
  na.rm.facet = TRUE,
  na.rm.var = TRUE,
  probs = NULL,
  prop_method = "beta",
  reorder = FALSE,

```

```

show_ci = TRUE,
show_n = FALSE,
show_value = TRUE,
show_labs = TRUE,
scale = 100,
digits = 0,
unit = "%",
dec = NULL,
col = "sienna2",
pal = NULL,
dodge = 0.9,
font = "Roboto",
wrap_width_y = 25,
title = NULL,
subtitle = NULL,
xlab = NULL,
ylab = NULL,
caption = NULL,
lang = "fr",
theme = "fonctionr",
coef_font = 1,
export_path = NULL
)

```

```
distrib_d(...)
```

Arguments

data	A dataframe or an object from the survey package or an object from the srvyr package.
quali_var	The discrete variable to be described.
facet	A variable defining the faceting group.
filter_exp	An expression filtering the data, preserving the design. Notice that <code>filter_exp</code> works as <code>srvyr::filter()</code> : it excludes observations for which <code>filter_exp</code> results into NA. It is often the case when NA is present on one of the filter variables.
...	All options possible in <code>srvyr::as_survey_design()</code> .
na.rm.facet	TRUE if you want to remove observations with NA on the facet variable. FALSE if you want to create a facet with the NA values for the facet variable. Default is TRUE.
na.rm.var	TRUE if you want to remove observations with NA on the discrete variable. FALSE if you want to create a modality with NA values for the discrete variable. Default is TRUE.
probs	Vector of probabilities for H0 of the statistical test, in the correct order (will be rescaled to sum to 1). If <code>probs = NULL</code> , no statistical test is performed. Default is NULL.
prop_method	Type of proportion method used to compute confidence intervals. See <code>survey::svyciprop()</code> for details. Default is beta method.

reorder	TRUE if you want to reorder the groups according to the proportion. NA value, if <code>na.rm.var = FALSE</code> , is not included in the reorder. In case of facets, the categories are reordered based on each median category. Default is FALSE.
show_ci	TRUE if you want to show the error bars on the graphic. FALSE if you don't want to show the error bars. Default is TRUE.
show_n	TRUE if you want to show on the graphic the number of observations in the sample in each category. FALSE if you don't want to show this number. Default is FALSE.
show_value	TRUE if you want to show the proportions in each category on the graphic. FALSE if you don't want to show the proportion. Default is TRUE.
show_labs	TRUE if you want to show axes labels. FALSE if you do not want to show any label on axes. Default is TRUE.
scale	Denominator of the proportion. Default is 100 to interpret numbers as percentages.
digits	Number of decimal places displayed on the values labels on the graphic. Default is 0.
unit	Unit displayed on the graphic. Default is "%".
dec	Decimal mark displayed on the graphic. Default depends on lang: ", " for fr and nl ; "." for en.
col	Color of the bars. col must be a R color or an hexadecimal color code. Default is "sienna2". The color of NA category (in case of <code>na.rm.var = FALSE</code>) is always "grey".
pal	Argument kept for compatibility with old versions.
dodge	Width of the bars. Default is 0.9 to let a small space between bars. A value of 1 leads to no space between bars. Values higher than 1 are not advised because they cause an overlapping of the bars.
font	Font used in the graphic. See <code>load_and_active_fonts()</code> for available fonts. Default is "Roboto".
wrap_width_y	Number of characters before going to the line for the labels of the categories. Default is 25.
title	Title of the graphic.
subtitle	Subtitle of the graphic.
xlab	X label on the graphic. As <code>ggplot2::coord_flip()</code> is used in the graphic, xlab refers to the x label on the graphic, after the <code>ggplot2::coord_flip()</code> , and not to the x variable in the data. Default (<code>xlab = NULL</code>) displays "Distribution (total : 100 pourcent)" (if lang = "fr"), "Distribution (total: 100 percent)" (if lang = "en") or "Distributie (totaal : 100 procent)" (if lang = "nl"). To show no X label, use <code>xlab = ""</code> .
ylab	Y label on the graphic. As <code>ggplot2::coord_flip()</code> is used in the graphic, ylab refers to the Y label on the graphic, after the <code>ggplot2::coord_flip()</code> , and not to the Y variable in the data. Default (<code>ylab = NULL</code>) displays the name of the discrete variable (<code>quali_var</code>). To show no Y label, use <code>ylab = ""</code> .
caption	Caption of the graphic. This caption goes under the default caption showing the result of the statistical test (if any).

lang	Language of the indications on the graphic. Possibilities are "fr" (french), "nl" (dutch) and "en" (english). Default is "fr".
theme	Theme of the graphic. Default is "fonctionr". "IWEPS" adds y axis lines and ticks. NULL uses the default grey ggplot2 theme.
coef_font	A multiplier factor for font size of all fonts on the graphic. Default is 1. Usefull when exporting the graphic for a publication (e.g. in a Quarto document).
export_path	Path to export the results in an xlsx file. The file includes two or three sheets : the table, the graphic and the statistical test (if probs is not NULL).

Value

A list that contains a table, a ggplot graphic and, if probs is not NULL, a statistical test.

Examples

```
# Loading of data
data(eusilc, package = "laeken")

# Recoding eusilc$pl030 into eusilc$pl030_rec
eusilc$pl030_rec <- NA
eusilc$pl030_rec[eusilc$pl030 == "1"] <- "Working full time"
eusilc$pl030_rec[eusilc$pl030 == "2"] <- "Working part time"
eusilc$pl030_rec[eusilc$pl030 == "3"] <- "Unemployed"
eusilc$pl030_rec[eusilc$pl030 == "4"] <- "Student"
eusilc$pl030_rec[eusilc$pl030 == "5"] <- "Retired"
eusilc$pl030_rec[eusilc$pl030 == "6"] <- "Permanently disabled"
eusilc$pl030_rec[eusilc$pl030 == "7"] <- "Fulfilling domestic tasks"

# Computation, taking sample design into account
eusilc_dist_group_d <- distrib_d(
  eusilc,
  pl030_rec,
  strata = db040,
  ids = db030,
  weight = rb050,
  title = "Distribution of socio-economic status",
  subtitle = "Example with austrian SILC data from 'laeken' package"
)

# Results in graph form
eusilc_dist_group_d$graph

# Results in table format
eusilc_dist_group_d$tab
```

`distrib_group_continuous`*distrib_group_continuous*

Description

Function to compare the distribution of a continuous variable between groups from complex survey data. It produces a list containing a density table (`dens`), a central value table (`tab`), a quantile table (`quant`), a ready-to-be published ggplot graphic (`graph`), a box-plot table (`moustache`) and a statistical test (`test`).

The density table contains x-y coordinates to draw density curve for each group. The central value table contains, for each group, the median or the mean of the continuous variable, with their confidence intervals, the sample size and the estimations of the totals, with their confidence intervals. The quantile table contains, for each group, quantiles and their confidence intervals. The box-plot table contains the X coordinates to draw the moustache, for each group.

In case of mean comparison, the statistical test is a Wald test (using `survey::regTermTest()`). In case of median comparison the statistical test is a Kruskal Wallis test (using `survey::svyranktest()`). The confidence intervals are taking into account the complex survey design.

Exporting those results to an Excell file is possible.

Usage

```
distrib_group_continuous(  
  data,  
  group,  
  quanti_exp,  
  type = "median",  
  facet = NULL,  
  filter_exp = NULL,  
  ...,  
  na.rm.group = TRUE,  
  na.rm.facet = TRUE,  
  quantiles = seq(0.1, 0.9, 0.1),  
  moustache_probs = c(0.95, 0.8, 0.5),  
  bw = 1,  
  resolution = 512,  
  height = 0.8,  
  limits = NULL,  
  reorder = FALSE,  
  show_mid_point = TRUE,  
  show_mid_line = FALSE,  
  show_ci_errorbar = TRUE,  
  show_ci_lines = FALSE,  
  show_ci_area = FALSE,  
  show_quant_lines = FALSE,  
  show_moustache = TRUE,
```

```

    show_value = TRUE,
    show_labs = TRUE,
    digits = 0,
    unit = "",
    dec = NULL,
    pal = NULL,
    col_density = "#e0dfe0",
    pal_moustache = NULL,
    col_moustache = c("#EB9BA0", "#FAD7B1"),
    color = NULL,
    col_border = NA,
    alpha = 1,
    font = "Roboto",
    wrap_width_y = 25,
    title = NULL,
    subtitle = NULL,
    xlab = NULL,
    ylab = NULL,
    caption = NULL,
    lang = "fr",
    theme = "fonctionr",
    coef_font = 1,
    export_path = NULL
  )

distrib_group_c(...)

```

Arguments

data	A dataframe or an object from the survey package or an object from the srvyr package.
group	A variable defining groups to be compared.
quanti_exp	An expression defining the quantitative variable the variable to be described and compared between groups. Notice that any observations with NA in at least one of the variable in <code>quanti_exp</code> are excluded for the computation of the densities and of the indicators.
type	Type of central value : "mean" to compute mean as the central value by group ; "median" to compute median as the central value by group.
facet	Not yet implemented.
filter_exp	An expression filtering the data, preserving the design. Notice that <code>filter_exp</code> works as <code>srvyr::filter()</code> : it excludes observations for which <code>filter_exp</code> results into NA. It is often the case when NA is present on one of the filter variables.
...	All options possible in <code>srvyr::as_survey_design()</code> .
na.rm.group	TRUE if you want to remove observations with NA on the group variable. FALSE if you want to create a group with the NA values for the group variable. Default is TRUE.

<code>na.rm.facet</code>	Not yet implemented.
<code>quantiles</code>	Quantiles computed in the distributions. Default are deciles.
<code>moustache_probs</code>	A vector defining the proportions of the population used to draw the boxplot. Default is <code>c(0.95, 0.8, 0.5)</code> to draw a boxplot with three groups containing respectively 50 percent, 80 percent and 95 percent of the population around to the median.
<code>bw</code>	The smoothing bandwidth to be used. The kernels are scaled such that this is the standard deviation of the smoothing kernel. Default is 1.
<code>resolution</code>	Resolution of the density curve. Default is 512.
<code>height</code>	Height of the curves. Default is 0.8. Values higher than 1 may cause curves to overlap.
<code>limits</code>	Limits of the x axe of the graphic. Does not apply to the computation. Default is NULL to show the entire distribution on the graphic. If the limits are shorter than the boxplot, some part of some boxplot will not be drawn.
<code>reorder</code>	TRUE if you want to reorder the groups according to the mean/median (depending on type). Unlike other functions, NA values, if <code>na.rm.group = FALSE</code> , is included in the reorder.
<code>show_mid_point</code>	TRUE if you want to show the mean or median (depending on type) as a point on the graphic. FALSE if you do not want to. Default is TRUE.
<code>show_mid_line</code>	TRUE if you want to show the mean or median (depending on type) as a line on the graphic. FALSE if you do not want to. Default is FALSE.
<code>show_ci_errorbar</code>	TRUE if you want to show confidence interval of the mean or median (depending on type) as an error bar on the graphic. FALSE if you do not want to show it as lines. Default is TRUE.
<code>show_ci_lines</code>	TRUE if you want to show confidence interval of the mean or median (depending on type) as lines on the graphic. FALSE if you do not want to show it as lines. Default is FALSE.
<code>show_ci_area</code>	TRUE if you want to show confidence interval of the mean or median (depending on type) as a coloured area on the graphic. FALSE if you do not want to show it as an area. Default is FALSE.
<code>show_quant_lines</code>	TRUE if you want to show quantiles as lines on the graphic. FALSE if you do not want to show them as lines. Default is FALSE.
<code>show_moustache</code>	TRUE if you want to show the boxplot on the graphic. FALSE if you do not want to show it. Default is TRUE.
<code>show_value</code>	TRUE if you want to show the value of mean/median of each group on the graphic. FALSE if you do not want to show the mean/median. Default is TRUE.
<code>show_labs</code>	TRUE if you want to show axes labels. FALSE if you do not want to show any labels on axes. Default is TRUE.
<code>digits</code>	Number of decimal places displayed on the values labels on the graphic. Default is 0.
<code>unit</code>	Unit displayed on the graphic. Default is none ("").

dec	Decimal mark shown on the graphic. Depends on lang: ", " for fr and nl ; "." for en.
pal	For compatibility with older versions.
col_density	Color of the density area. It may be one color or a vector with several colors. Colors should be R color or an hexadecimal color code. In case of one color, the density is monocolour. In case of a vector, the quantile areas are painted in continuous colors going from the last color in the vector (center quantile) to the first color (first and last quantiles). In case of an even quantile area numbers (e.g. deciles, quartiles) the last color of the vector is only applied to the highcenter quantile area to avoid two continuous quantile areas having the same color.
pal_moustache	For compatibility with old versions.
col_moustache	Color of the moustache. Can be one or several colors to create a palette. In case of a vector, the different areas of the box-plot are painted in continuous colors going from the first color in the vector (center of the box-plot) to the last color (extern area of the box-plot).
color	For compatibility with older versions.
col_border	Color of the density line. Color should be one R color or one hexadecimal color code. Default (NULL) does not draw the density line.
alpha	Transparence of the density areas. Default is 1. It applies only to col_density.
font	Font used in the graphic. See load_and_active_fonts() for available fonts. Default is "Roboto".
wrap_width_y	Number of characters before going to the line for the labels of the groups. Default is 25.
title	Title of the graphic.
subtitle	Subtitle of the graphic.
xlab	X label on the graphic. If xlab = NULL, X label on the graphic will be quanti_exp.
ylab	Y label on the graphic. Default (ylab = NULL) displays the name of the group variable. To show no Y label, use ylab = "".
caption	Caption of the graphic. This caption goes under the default caption showing the result of the Chi-Square test. There is no way of not showing the result of the statistical test as a caption.
lang	Language of the indications on the graphic. Possibilities are "fr" (french), "nl" (dutch) and "en" (english). Default is "fr".
theme	Theme of the graphic. Default is "fonctionr". "IWEPS" adds y axis lines and ticks. NULL uses the default grey ggplot2 theme.
coef_font	A multiplier factor for font size of all fonts on the graphic. Default is 1. Usefull when exporting the graphic for a publication (e.g. in a Quarto document).
export_path	Path to export the results in an xlsx file. The file includes five sheets: the central values table, the quantile table, the densities table, the graphic and the statistical test result.

Value

A list that contains a density table (dens), a central values table (tab), a quantile table (quant), a ggplot graphic (graph), boxplot table (moustache) and a statistical test (test).

Examples

```

# Loading of data
data(eusilc, package = "laeken")

# Recoding eusilc$pl030 into eusilc$pl030_rec
eusilc$pl030_rec <- NA
eusilc$pl030_rec[eusilc$pl030 == "1"] <- "Working full time"
eusilc$pl030_rec[eusilc$pl030 == "2"] <- "Working part time"
eusilc$pl030_rec[eusilc$pl030 == "3"] <- "Unemployed"
eusilc$pl030_rec[eusilc$pl030 == "4"] <- "Student"
eusilc$pl030_rec[eusilc$pl030 == "5"] <- "Retired"
eusilc$pl030_rec[eusilc$pl030 == "6"] <- "Permanently disabled"
eusilc$pl030_rec[eusilc$pl030 == "7"] <- "Fulfilling domestic tasks"

# Computation, taking sample design into account
eusilc_dist_g_c <- distrib_group_c(
  eusilc,
  group = pl030_rec,
  quanti_exp = eqIncome,
  strata = db040,
  ids = db030,
  weight = rb050,
  limits = c(0, 50000),
  resolution = 128,
  title = "Distribution of eq. income",
  subtitle = "Example with austrian SILC data from 'laeken' package"
)

# Results in graph form
eusilc_dist_g_c$graph

# Results in table format
eusilc_dist_g_c$tab

```

distrib_group_discrete

distrib_group_discrete

Description

Function to compare the distribution of a discrete variable between different groups based on complex survey data.

It produces a list containing a table, including the confidence intervals of the indicators, a ready-to-be published ggplot graphic and a Chi-Square statistical test (using `survey::svychisq()`). The confidence intervals and the statistical test are taking into account the complex survey design. In case of facets, no statistical test is (yet) computed.

Exporting those results to an Excell file is possible.

Usage

```
distrib_group_discrete(  
  data,  
  group,  
  quali_var,  
  facet = NULL,  
  filter_exp = NULL,  
  ...,  
  na.rm.group = TRUE,  
  na.rm.facet = TRUE,  
  na.rm.var = TRUE,  
  total = TRUE,  
  prop_method = "beta",  
  reorder = FALSE,  
  show_n = FALSE,  
  show_value = TRUE,  
  show_labs = TRUE,  
  total_name = NULL,  
  scale = 100,  
  digits = 0,  
  unit = "",  
  dec = NULL,  
  pal = "OBSS",  
  direction = 1,  
  desaturate = 0,  
  lighten = 0,  
  darken = 0,  
  dodge = 0.9,  
  font = "Roboto",  
  wrap_width_y = 25,  
  wrap_width_leg = 25,  
  legend_ncol = 4,  
  title = NULL,  
  subtitle = NULL,  
  xlab = NULL,  
  ylab = NULL,  
  legend_lab = NULL,  
  caption = NULL,  
  lang = "fr",  
  theme = "fonctionr",  
  coef_font = 1,  
  export_path = NULL  
)  
  
distrib_group_d(...)
```

Arguments

<code>data</code>	A dataframe or an object from the survey package or an object from the srvyr package.
<code>group</code>	A variable defining groups to be compared.
<code>quali_var</code>	The discrete variable described among the different groups.
<code>facet</code>	A variable defining the faceting group.
<code>filter_exp</code>	An expression filtering the data, preserving the design. Notice that <code>filter_exp</code> works as <code>srvyr::filter()</code> : it excludes observations for which <code>filter_exp</code> results into NA. It is often the case when NA is present on one of the filter variables.
<code>...</code>	All options possible in <code>srvyr::as_survey_design()</code> .
<code>na.rm.group</code>	TRUE if you want to remove observations with NA on the group. FALSE if you want to create a group with the NA values for the group variable. Default is TRUE.
<code>na.rm.facet</code>	TRUE if you want to remove observations with NA on the facet variable. FALSE if you want to create a facet with the NA values for the facet variable. Default is TRUE.
<code>na.rm.var</code>	TRUE if you want to remove observations with NA on the discrete variable. FALSE if you want to create a modality with NA values for the discrete variable. Default is TRUE.
<code>total</code>	TRUE if you want to compute a total, FALSE if you don't. The default is TRUE.
<code>prop_method</code>	Type of proportion method used to compute confidence intervals. See <code>survey::svyciprop()</code> for details. Default is beta method.
<code>reorder</code>	TRUE if you want to reorder the groups according to the proportion of the first level of <code>quali_var</code> . NA group, if <code>na.rm.group = FALSE</code> , is not included in the reorder. In case of facets, the groups are reordered based on each median group. Default is FALSE.
<code>show_n</code>	TRUE if you want to show on the graphic the number of observations in the sample in each category (of <code>quali_var</code>) of each group. FALSE if you don't want to show this number. Default is FALSE.
<code>show_value</code>	TRUE if you want to show the proportion in each category of each group on the graphic. FALSE if you do not want to show the proportions. Proportions of 2 percent or less are never showed on the graphic. Default is TRUE.
<code>show_labs</code>	TRUE if you want to show axes and legend labels. FALSE if you don't want to show any labels on axes and legend. Default is TRUE.
<code>total_name</code>	Name of the total displayed on the graphic. Default is "Total" in French and in English and "Totaal" in Dutch.
<code>scale</code>	Denominator of the proportions. Default is 100 to interpret numbers as percentages.
<code>digits</code>	Number of decimal places displayed on the values labels on the graphic. Default is 0.
<code>unit</code>	Unit showed in the graphic. Default (<code>unit = ""</code>) shows not unit on values and percent on the X axe.

dec	Decimal mark shown on the graphic. Depends on lang: ", " for fr and nl ; "." for en.
pal	Colors of the bars. pal must be vector of R colors or hexadecimal colors or a palette from packages MetBrewer or PrettyCols or a palette from fonctionr. The color of NA category (in case of na.rm.var = FALSE) is always "grey".
direction	Direction of the palette color. Default is 1. The opposite direction is -1.
desaturate	Numeric specifying the amount of desaturation where 1 corresponds to complete desaturation (no colors, grey layers only), 0 to no desaturation, and values in between to partial desaturation. Default is 0. See colorspace::desaturate() for details. If desaturate and lighten/darken arguments are used, lighten/darken is applied in a second time (i.e. on the color transformed by desaturate).
lighten	Numeric specifying the amount of lightening. Negative numbers cause darkening. Value should be ranged between -1 (black) and 1 (white). Default is 0. It doesn't affect the color of NA (in case of na.rm.group = FALSE). See colorspace::lighten() for details. If both argument lighten and darken are used (not advised), darken is applied in a second time (i.e. on the color transformed by lighten).
darken	Numeric specifying the amount of lightening. Negative numbers cause lightening. Value should be ranged between -1 (white) and 1 (black). Default is 0. It doesn't affect the color of NA (in case of na.rm.group = FALSE). See colorspace::darken() for details. If both argument lighten and darken are used (not advised), darken is applied in a second time (i.e. on the color transformed by lighten).
dodge	Width of the bars. Default is 0.9 to let a small space between bars. A value of 1 leads to no space between bars. Values higher than 1 are not advised because they cause an overlapping of the bars.
font	Font used in the graphic. See load_and_active_fonts() for available fonts. Default is "Roboto".
wrap_width_y	Number of characters before going to the line for the labels of the groups. Default is 25.
wrap_width_leg	Number of characters before going to the line for the labels of quali_var. Default is 25.
legend_ncol	Number of columns in the legend. Default is 4.
title	Title of the graphic.
subtitle	Subtitle of the graphic.
xlab	X label on the graphic. As ggplot2::coord_flip() is used in the graphic, xlab refers to the x label on the graphic, after the ggplot2::coord_flip() , and not to the x variable in the data. Default (xlab = NULL) displays "Distribution : " (if lang = "fr"), "Distribution: " (if lang = "en") or "Distributie: " (if lang = "nl"), followed by the name of the discrete variable (quali_var). To show no X label, use xlab = "".
ylab	Y label on the graphic. As ggplot2::coord_flip() is used in the graphic, ylab refers to the y label on the graphic, after the ggplot2::coord_flip() , and not to the y variable in the data. Default (ylab = NULL) displays the name of the group variable. To show no Y label, use ylab = "".

legend_lab	Legend (fill) label on the graphic. Default (legend_lab = NULL) displays the name of the discrete variable (quali_var). To show no legend label, use legend_lab = "".
caption	Caption of the graphic. This caption goes under the default caption showing the result of the Chi-Square test. There is no way of not showing the result of the chi-square test as a caption.
lang	Language of the indications on the graphic. Possibilities are "fr" (french), "nl" (dutch) and "en" (english). Default is "fr".
theme	Theme of the graphic. Default is "fonctionr". "IWEPS" adds y axis lines and ticks. NULL uses the default grey ggplot2 theme.
coef_font	A multiplier factor for font size of all fonts on the graphic. Default is 1. Useful when exporting the graphic for a publication (e.g. in a Quarto document).
export_path	Path to export the results in an xlsx file. The file includes three (without facets) or two sheets (with facets): the table, the graphic and the Chi-Square statistical test result.

Value

A list that contains a table, a ggplot graphic and, in most cases, a Chi-square statistical test.

Examples

```
# Loading of data
data(eusilc, package = "laeken")

# Recoding eusilc$pl030 into eusilc$pl030_rec
eusilc$pl030_rec <- NA
eusilc$pl030_rec[eusilc$pl030 == "1"] <- "Working full time"
eusilc$pl030_rec[eusilc$pl030 == "2"] <- "Working part time"
eusilc$pl030_rec[eusilc$pl030 == "3"] <- "Unemployed"
eusilc$pl030_rec[eusilc$pl030 == "4"] <- "Student"
eusilc$pl030_rec[eusilc$pl030 == "5"] <- "Retired"
eusilc$pl030_rec[eusilc$pl030 == "6"] <- "Permanently disabled"
eusilc$pl030_rec[eusilc$pl030 == "7"] <- "Fulfilling domestic tasks"

# Computation, taking sample design into account
eusilc_dist_d <- distrib_group_d(
  eusilc,
  group = pb220a,
  quali_var = pl030_rec,
  strata = db040,
  ids = db030,
  weight = rb050,
  title = "Distribution of socio-economic status according to nationality",
  subtitle = "Example with austrian SILC data from 'laeken' package"
)

# Results in graph form
eusilc_dist_d$graph
```

```
# Results in table format
eusilc_dist_d$tab
```

esth_graph	<i>esth_graph</i>
------------	-------------------

Description

Function to construct a graphic following the aesthetics of the other functions of `functionr` from a table. This function was created to align results generated outside `fonctionr` with the outputs of `fonctionr`.

Usage

```
esth_graph(  
  tab,  
  var,  
  value,  
  error_low = NULL,  
  error_upp = NULL,  
  facet = NULL,  
  n_var = NULL,  
  pvalue = NULL,  
  reorder = FALSE,  
  show_value = TRUE,  
  name_total = NULL,  
  scale = 1,  
  digits = 2,  
  unit = "",  
  dec = ",",  
  pal = NULL,  
  col = "indianred4",  
  dodge = 0.9,  
  font = "Roboto",  
  wrap_width_y = 25,  
  title = NULL,  
  subtitle = NULL,  
  xlab = NULL,  
  ylab = NULL,  
  caption = NULL,  
  theme = "fonctionr",  
  coef_font = 1  
)
```

Arguments

`tab` dataframe with the indicators to be plotted.

var	The variable in tab with the labels of the indicator to be plotted.
value	The variable in tab with the values of the indicator to be plotted.
error_low	The variable in tab with the lower bound of the confidence interval. If either error_low or error_upp is NULL error bars are not shown on the graphic.
error_upp	The variable in tab with the upper bound of the confidence interval. If either error_low or error_upp is NULL error bars are not shown on the graphic.
facet	A variable in tab defining the faceting group, if applicable. Default is NULL.
n_var	The variable in tab containing the number of observations for each indicator plotted. Default (NULL) does not show the numbers of observations on the plot.
pvalue	The p-value to show in the caption. It can be a numeric value or the pvalue object from a statistical test.
reorder	TRUE if you want to reorder var according to value. FALSE if you do not want to reorder. NA and total labels in var are not included in the reorder. Default is FALSE.
show_value	TRUE if you want to show the values on the graphic. FALSE if you do not want to show them. Default is TRUE.
name_total	Name of the var label that may contain the total. When indicated, it is displayed separately (bold name and value color is 'grey40') on the graph.
scale	Denominator of the indicator. Default is 1 to not modify indicators.
digits	Number of decimal places displayed on the values labels on the graphic. Default is 0.
unit	The unit displayed on the graphic. Default is no unit ("").
dec	Decimal mark shown on the graphic. Default is ", ".
pal	For compatibility with old versions.
col	Color of the bars. col must be a R color or an hexadecimal color code. Default is "indianred4". The color of NA and total are always "grey" and "grey40".
dodge	Width of the bars. Default is 0.9 to let a small space between bars. A value of 1 leads to no space between bars. Values higher than 1 are not advised because they cause an overlapping of the bars.
font	Font used in the graphic. See load_and_active_fonts() for available fonts. Default is "Roboto".
wrap_width_y	Number of characters before going to the line for the labels of var Default is 25.
title	Title of the graphic.
subtitle	Subtitle of the graphic.
xlab	X label on the graphic. As ggplot2::coord_flip() is used in the graphic, xlab refers to the x label on the graphic, after the ggplot2::coord_flip(), and not to var in tab.
ylab	Y label on the graphic. As ggplot2::coord_flip() is used in the graphic, ylab refers to the y label on the graphic, after the ggplot2::coord_flip(), and not to value in tab.
caption	Caption of the graphic.

theme	Theme of the graphic. Default is "fonctionr". "IWEPS" adds y axis lines and ticks. NULL uses the default grey ggplot2 theme.
coef_font	A multiplier factor for font size of all fonts on the graphic. Default is 1. Usefull when exporting the graphic for a publication (e.g. in a Quarto document).

Value

A ggplot graphic.

Examples

```
# Making fictional dataframe
data_test <- data.frame(
  Indicators = c(
    "Variable 1",
    "Variable 2",
    "Variable 3",
    "Variable 4",
    "Variable 5",
    "Tot"
  ),
  Estimates = c(1.52, 1.63, 2.34, 4.15, 1.32, 2.13),
  IC_low = c(1.32, 1.4, 1.98, 4, 14.2, 26),
  IC_upp = c(1.73, 1.81, 22.4, 47.44, 1.45, 2.34),
  sample_size = c(215, 300, 129, 212, 189, 1045)
)

# Using dataframe to make a plot
plot_test <- esth_graph(data_test,
  var = Indicators,
  value = Estimates,
  error_low = IC_low,
  error_upp = IC_upp,
  n_var = sample_size,
  pvalue = .001,
  reorder = TRUE,
  show_value = TRUE,
  name_total = "Tot",
  scale = 1,
  digits = 1,
  unit = "%",
  dec = ".",
  col = "green4",
  dodge = 0.8,
  font = "Montserrat",
  wrap_width_y = 25,
  title = "Plot",
  subtitle = "Using fake data",
  xlab = "Proportion (in %)",
  ylab = "Indicators",
  caption = "Source: fictional own calculation",
  theme = "IWEPS"
```

```
)  
  
# Result is a ggplot  
plot_test
```

fonctionr_options *fonctionr_options*

Description

Function to set global options for fonctionr. The arguments defined in the options are only active if the user has not manually specified a value for those arguments within the various functions. Arguments may be shared by multiple functions (if they have the same name) or specific to certain functions.

Usage

```
fonctionr_options(  
  na.rm.group = NULL,  
  na.rm.facet = NULL,  
  na.prop = NULL,  
  na.vars = NULL,  
  na.rm.var = NULL,  
  probs = NULL,  
  total = NULL,  
  prop_method = NULL,  
  quantiles = NULL,  
  moustache_probs = NULL,  
  bw = NULL,  
  resolution = NULL,  
  height = NULL,  
  limits = NULL,  
  reorder = NULL,  
  position = NULL,  
  show_ci = NULL,  
  show_mid_point = NULL,  
  show_mid_line = NULL,  
  show_ci_errorbar = NULL,  
  show_ci_lines = NULL,  
  show_ci_area = NULL,  
  show_quant_lines = NULL,  
  show_moustache = NULL,  
  show_n = NULL,  
  show_value = NULL,  
  show_labs = NULL,  
  total_name = NULL,  
  scale = NULL,
```

```

digits = NULL,
unit = NULL,
dec = NULL,
col = NULL,
pal = NULL,
direction = NULL,
desaturate = NULL,
lighten = NULL,
darken = NULL,
col_density = NULL,
col_moustache = NULL,
col_border = NULL,
alpha = NULL,
dodge = NULL,
font = NULL,
wrap_width_y = NULL,
wrap_width_leg = NULL,
legend_ncol = NULL,
title = NULL,
subtitle = NULL,
xlab = NULL,
ylab = NULL,
legend_lab = NULL,
caption = NULL,
lang = NULL,
theme = NULL,
coef_font = NULL,
parallel = NULL,
erase_all = FALSE
)

```

Arguments

na.rm.group	na.rm.group argument.
na.rm.facet	na.rm.facet argument.
na.prop	na.prop argument.
na.vars	na.vars argument.
na.rm.var	na.rm.var argument.
probs	probs argument.
total	total argument.
prop_method	prop_method argument.
quantiles	quantiles argument.
moustache_probs	moustache_probs argument.
bw	bw argument.
resolution	resolution argument.

height	height argument.
limits	limits argument.
reorder	reorder argument.
position	position argument.
show_ci	show_ci argument.
show_mid_point	show_mid_point argument.
show_mid_line	show_mid_line argument.
show_ci_errorbar	show_ci_errorbar argument.
show_ci_lines	show_ci_lines argument.
show_ci_area	show_ci_area argument.
show_quant_lines	show_quant_lines argument.
show_moustache	show_moustache argument.
show_n	show_n argument.
show_value	show_value argument.
show_labs	show_labs argument.
total_name	total_name argument.
scale	scale argument.
digits	digits argument.
unit	unit argument.
dec	dec argument.
col	col argument.
pal	pal argument.
direction	direction argument.
desaturate	desaturate argument.
lighten	lighten argument.
darken	darken argument.
col_density	col_density argument.
col_moustache	col_moustache argument.
col_border	col_border argument.
alpha	alpha argument.
dodge	dodge argument.
font	font argument.
wrap_width_y	wrap_width_y argument.
wrap_width_leg	wrap_width_leg argument.
legend_ncol	legend_ncol argument.
title	title argument.

subtitle	subtitle argument.
xlab	xlab argument.
ylab	ylab argument.
legend_lab	legend_lab argument.
caption	caption argument.
lang	lang argument.
theme	theme argument.
coef_font	coef_font argument.
parallel	parallel argument.
erase_all	TRUE erases all the options. Default is FALSE.

Value

No return value, called for side effects.

Examples

```
# We set global settings
fonctionr_options(coef_font = 1.5, col = "magenta", caption = "Beautiful caption")

# Loading of data
data(eusilc, package = "laeken")

# Recoding eusilc$pl030 into eusilc$pl030_rec
eusilc$pl030_rec <- NA
eusilc$pl030_rec[eusilc$pl030 == "1"] <- "Working full time"
eusilc$pl030_rec[eusilc$pl030 == "2"] <- "Working part time"
eusilc$pl030_rec[eusilc$pl030 == "3"] <- "Unemployed"
eusilc$pl030_rec[eusilc$pl030 == "4"] <- "Student"
eusilc$pl030_rec[eusilc$pl030 == "5"] <- "Retired"
eusilc$pl030_rec[eusilc$pl030 == "6"] <- "Permanently disabled"
eusilc$pl030_rec[eusilc$pl030 == "7"] <- "Fulfilling domestic tasks"

# Computation, taking sample design into account
eusilc_prop <- prop_group(
  eusilc,
  group = pl030_rec,
  prop_exp = py090n > 0,
  weight = rb050,
  title = "% of ind. receiving unemployment benefits in their hh",
  subtitle = "Example with austrian SILC data from 'laeken' package"
)

# Results in graph form
eusilc_prop$graph

# We set back settings to default
fonctionr_options(erase_all = TRUE)
```

load_and_active_fonts *load_and_active_fonts*

Description

Function to load and activate fonctionr's built-in fonts. Available fonts, included in the package itself, are "Roboto", "Montserrat", "Gotham Narrow", and "Euclid Circular A". Default is "Roboto".

Usage

```
load_and_active_fonts()
```

Value

No return value, called for side effects.

Examples

```
# Loading of fonts from fonctionr. You can now use it in fonctionr !
load_and_active_fonts()
```

make_surface *make_surface*

Description

Function that represents the values of a group variable as areas proportional to those values.

Usage

```
make_surface(
  tab,
  var,
  value,
  error_low = NULL,
  error_upp = NULL,
  facet = NULL,
  pvalue = NULL,
  reorder = FALSE,
  compare = FALSE,
  space = NULL,
  position = "mid",
  show_ci = TRUE,
  name_total = "Total",
  digits = 0,
```

```

    unit = NULL,
    col = NULL,
    pal = "OBSS_Autumn",
    direction = 1,
    desaturate = 0,
    lighten = 0,
    darken = 0,
    size_text = 3.88,
    bg = "#f8f5f5",
    linewidth_ci = 0.5,
    ratio = 3/2,
    font = "Roboto",
    wrap_width_lab = 20,
    title = NULL,
    subtitle = NULL,
    hjust.title = 0,
    caption = NULL,
    coef_font = 1
  )

```

Arguments

tab	dataframe with the variables to be plotted.
var	The variable in tab with the labels of the indicator to be plotted.
value	The variable in tab with the values of the indicator to be plotted.
error_low	The variable in tab that is the lower bound of the confidence interval. If either error_low or error_upp is NULL error rectangles are not shown on the graphic.
error_upp	The variable in tab that is the upper bound of the confidence interval. If either error_low or error_upp is NULL error rectangles are not shown on the graphic.
facet	A variable in tab defining the faceting group, if applicable. Default is NULL.
pvalue	The p-value to show in the caption. It can be a numeric value or the pvalue object from a statistical test.
reorder	TRUE if you want to reorder the values. NA label in var is not included in the reorder.
compare	TRUE to display a rectangle representing the smallest value. When facets are enabled, this is the smallest value per facet category.
space	The space between the rectangles. The unit is that of the indicator.
position	The position of the rectangles: "mid" for center alignment, "bottom" for bottom alignment.
show_ci	TRUE if you want to show the CI on the graphic. The bounds of the confidence intervals are displayed as dotted rectangles around the result. FALSE if you do not want to show them. Default is TRUE.
name_total	Name of the var label that may contain the total. When indicated, it is not displayed on the graph.

digits	Number of decimal places displayed on the values labels on the graphic. Default is 0.
unit	The unit showd on the plot. Default is none ("").
col	Color of the rectangles if the user wants a monocolour graph. col must be a R color or an hexadecimal color code. As pal has a priority over col, if the user wants to use col, he must not use simultaneously the pal argument (even pal = NULL).
pal	Colors of the rectangles if the user wants the rectangles to have different colors. pal must be vector of R colors or hexadecimal colors or a palette from packages MetBrewer or PrettyCols or a palette from fonctionr. pal has a priority over col.
direction	Direction of the palette color. Default is 1. The opposite direction is -1.
desaturate	Numeric specifying the amount of desaturation where 1 corresponds to complete desaturation (no colors, grey layers only), 0 to no desaturation, and values in between to partial desaturation. Default is 0. It affects only the palette (pal) and not the monocolour (col). See colorspace::desaturate() for details. If desaturate and lighten/darken arguments are used, lighten/darken is applied in a second time (i.e. on the color transformed by desaturate).
lighten	Numeric specifying the amount of lightening. Negative numbers cause darkening. Value should be ranged between -1 (black) and 1 (white). Default is 0. It affects only the palette (pal) and not the monocolour (col). See colorspace::lighten() for details. If both argument lighten and darken are used (not advised), darken is applied in a second time (i.e. on the color transformed by lighten).
darken	Numeric specifying the amount of lightening. Negative numbers cause lightening. Value should be ranged between -1 (white) and 1 (black). Default is 0. It affects only the palette (pal) and not the monocolour (col). See colorspace::darken() for details. If both argument lighten and darken are used (not advised), darken is applied in a second time (i.e. on the color transformed by lighten).
size_text	Text size displayed in rectangles . Default is 3.88 (as in ggplot2).
bg	Color of the background. bg must be a R color or an hexadecimal color code.
linewidth_ci	Line width of the dotted confidence intervals lines. It affects also the lengths of the dots and spaces between dots. Default is 0.5 to have confidence lines two times thinner than the lines of the indicators.
ratio	Ratio between the length and the width of the rectangles. 1 produces squares ; greater than 1 produces vertical rectangles and smaller than 1 produces horizontal rectangles. Default is 3/2.
font	Font used in the graphic. See load_and_active_fonts() for available fonts. Default is "Roboto".
wrap_width_lab	Number of characters before going to the line for the labels of the categories of var. Default is 20.
title	Title of the graphic.
subtitle	Subtitle of the graphic.
hjust.title	Horizontal alignment of title & subtitle. It should take a numeric value. Default (0) leads to left alignment, 1 leads to right alignment and 0.5 leads to centered alignment.

caption	Caption of the graphic.
coef_font	A multiplier factor for font size of all fonts on the graphic. Default is 1. Useful when exporting the graphic for a publication (e.g. in a Quarto document).

Value

A ggplot graphic.

Examples

```
# Loading of data
data(eusilc, package = "laeken")

# Recoding eusilc$pl030 into eusilc$pl030_rec
eusilc$pl030_rec <- NA
eusilc$pl030_rec[eusilc$pl030 == "1"] <- "Working full time"
eusilc$pl030_rec[eusilc$pl030 == "2"] <- "Working part time"
eusilc$pl030_rec[eusilc$pl030 == "3"] <- "Unemployed"
eusilc$pl030_rec[eusilc$pl030 == "4"] <- "Student"
eusilc$pl030_rec[eusilc$pl030 == "5"] <- "Retired"
eusilc$pl030_rec[eusilc$pl030 == "6"] <- "Permanently disabled"
eusilc$pl030_rec[eusilc$pl030 == "7"] <- "Fulfilling domestic tasks"

# Calculation of income means by age category with fonctionr, taking sample design into account
eusilc_mean <- mean_group(
  eusilc,
  group = pl030_rec,
  quanti_exp = py010n + py050n + py090n + py100n + py110n + py120n + py130n + py140n,
  filter_exp = !pl030_rec %in% c("Student", "Fulfilling domestic tasks") & db040 == "Tyrol",
  weights = rb050
)

# Displaying results with make_surface()
eusilc_mean$tab |>
  make_surface(
    var = pl030_rec,
    value = mean,
    error_low = mean_low,
    error_upp = mean_upp,
    reorder = TRUE,
    wrap_width_lab = 15,
    unit = "€",
    title = "Equivalentised income in household by socio-economic status",
    subtitle = "Example with austrian SILC data from 'laeken' package"
  )
```

Description

Function to compute the proportions of a set of several binary variables or means or medians of a set of quantitative variables, based on complex survey data.

It produces a list containing a table, including the confidence intervals of the indicators and a ready-to-be published ggplot graphic. The confidence intervals are taking into account the complex survey design.

Exporting the results to an Excell file is possible.

Usage

```
many_val(  
  data,  
  list_vars,  
  type,  
  list_vars_lab = NULL,  
  facet = NULL,  
  filter_exp = NULL,  
  ...,  
  na.rm.facet = TRUE,  
  na.vars = "rm",  
  prop_method = "beta",  
  reorder = FALSE,  
  show_ci = TRUE,  
  show_n = FALSE,  
  show_value = TRUE,  
  show_labs = TRUE,  
  scale = NULL,  
  digits = 0,  
  unit = NULL,  
  dec = NULL,  
  col = NULL,  
  pal = "OBSS_alt3",  
  direction = 1,  
  desaturate = 0,  
  lighten = 0,  
  darken = 0,  
  dodge = 0.9,  
  font = "Roboto",  
  wrap_width_y = 25,  
  title = NULL,  
  subtitle = NULL,  
  xlab = NULL,  
  ylab = NULL,  
  caption = NULL,  
  lang = "fr",  
  theme = "fonctionr",  
  coef_font = 1,  
  export_path = NULL
```

```

)

many_prop(..., type = "prop")

many_median(..., type = "median")

many_mean(..., type = "mean")

```

Arguments

<code>data</code>	A dataframe or an object from the survey package or an object from the srvyr package.
<code>list_vars</code>	A vector containing the names of the dummy/quantitative variables on which to compute the proportions/means/medians.
<code>type</code>	"prop" to compute proportions ; "mean" to compute means ; "median" to compute medians.
<code>list_vars_lab</code>	A vector containing the labels of the dummy/quantitative variables to be displayed on the graphic and in the table of result. Default uses the variable names in <code>list_vars</code> .
<code>facet</code>	A variable defining the faceting group.
<code>filter_exp</code>	An expression filtering the data, preserving the design. Notice that <code>filter_exp</code> works as <code>srvyr::filter()</code> : it excludes observations for which <code>filter_exp</code> results into NA. It is often the case when NA is present on one of the filter variables.
<code>...</code>	All options possible in <code>srvyr::as_survey_design()</code> .
<code>na.rm.facet</code>	TRUE if you want to remove observations with NA on the facet variable. FALSE if you want to create a facet with the NA values for the facet variable. Default is TRUE.
<code>na.vars</code>	The treatment of NA values in variables (<code>list_vars</code>). "rm" removes NA separately in each individual variable, "rm.all" removes every individual that has at least one NA in one variable. Default is "rm".
<code>prop_method</code>	Type of proportion method used to compute confidence intervals. See <code>survey::svyciprop()</code> for details. Default is beta method. This argument is only used in case of type = "prop".
<code>reorder</code>	TRUE if you want to reorder the variables according to the proportions/means/medians. Default is FALSE.
<code>show_ci</code>	TRUE if you want to show the error bars on the graphic. FALSE if you don't want to show the error bars. Default is TRUE.
<code>show_n</code>	TRUE if you want to show on the graphic the number of observations in the sample for each variable. The number can varie if <code>na.vars = "rm"</code> . FALSE if you do not want to show this number. Default is FALSE.
<code>show_value</code>	TRUE if you want to show the proportions/means/median for each variable on the graphic. FALSE if you do not want to show the proportions/means/medians. Default is TRUE.
<code>show_labs</code>	TRUE if you want to show axes labels. FALSE if you do not want to show any labels on axes. Default is TRUE.

scale	Denominator of the proportions. Default is 100 to interpret numbers as percentages. This argument is only used in case of type = "prop".
digits	Number of decimal places displayed on the values labels on the graphic. Default is 0.
unit	Unit displayed on the graphic. Default is percent for type = "prop" and no unit for type = "mean" or "median".
dec	Decimal mark displayed on the graphic. Default depends on lang: ", " for fr and nl ; "." for en.
col	Color of the bars if the user wants a monocolour graph. col must be a R color or an hexadecimal color code. As pal has a priority over col, if the user wants to use col, he must not use simultaneously the pal argument (even pal = NULL).
pal	Colors of the bars if the user wants the bars to have different colors. pal must be vector of R colors or hexadecimal colors or a palette from packages MetBrewer or PrettyCols or a palette from fonctionr. pal has a priority over col.
direction	Direction of the palette color. Default is 1. The opposite direction is -1.
desaturate	Numeric specifying the amount of desaturation where 1 corresponds to complete desaturation (no colors, grey layers only), 0 to no desaturation, and values in between to partial desaturation. Default is 0. It affects only the palette (pal) and not the monocolour (col). See colorspace::desaturate() for details. If desaturate and lighten/darken arguments are used, lighten/darken is applied in a second time (i.e. on the color transformed by desaturate).
lighten	Numeric specifying the amount of lightening. Negative numbers cause darkening. Value should be ranged between -1 (black) and 1 (white). Default is 0. It affects only the palette (pal) and not the monocolour (col). See colorspace::lighten() for details. If both argument lighten and darken are used (not advised), darken is applied in a second time (i.e. on the color transformed by lighten).
darken	Numeric specifying the amount of lightening. Negative numbers cause lightening. Value should be ranged between -1 (white) and 1 (black). Default is 0. It affects only the palette (pal) and not the monocolour (col). See colorspace::darken() for details. If both argument lighten and darken are used (not advised), darken is applied in a second time (i.e. on the color transformed by lighten).
dodge	Width of the bars. Default is 0.9 to let a small space between bars. A value of 1 leads to no space between bars. Values higher than 1 are not advised because they cause an overlapping of the bars.
font	Font used in the graphic. See load_and_active_fonts() for available fonts. Default is "Roboto".
wrap_width_y	Number of characters before going to the line for the labels of the groups. Default is 25.
title	Title of the graphic.
subtitle	Subtitle of the graphic.
xlab	X label on the graphic. As ggplot2::coord_flip() is used in the graphic, xlab refers to the x label on the graphic, after the ggplot2::coord_flip() , and not to the x variable in the data. Default (xlab = NULL) displays, for type =

	prop, "Proportion :." (if lang = "fr"), "Proportion:" (if lang = "en") or "Aandeel:" (if lang = "nl"), or, for type = "mean", "Moyenne :." (if lang = "fr"), "Mean:" (if lang = "en") or "Gemiddelde:" (if lang = "nl"), or, for type = "median", "Médiane :." (if lang = "fr"), "Median:" (if lang = "en") or "Medi-aan:" (if lang = "nl"), followed by the labels of the variables (list_vars_lab). To show no X label, use xlab = "".
ylab	Y label on the graphic. As <code>ggplot2::coord_flip()</code> is used in the graphic, ylab refers to the y label on the graphic, after the <code>ggplot2::coord_flip()</code> , and not to the y variable in the data. Default (ylab = NULL) displays no Y label.
caption	Caption of the graphic.
lang	Language of the indications on the graphic. Possibilities are "fr" (french), "nl" (dutch) and "en" (english). Default is "fr".
theme	Theme of the graphic. Default is "fonctionr". "IWEPS" adds y axis lines and ticks. NULL uses the default grey ggplot2 theme.
coef_font	A multiplier factor for font size of all fonts on the graphic. Default is 1. Usefull when exporting the graphic for a publication (e.g. in a Quarto document).
export_path	Path to export the results in an xlsx file. The file includes two sheets: the table and the graphic.

Value

A list that contains a table and a ggplot graphic.

Examples

```
# Loading of data
data(eusilc, package = "laeken")

# Recoding variables
eusilc$worker <- 0
eusilc$worker[eusilc$pl030 == "1"]<-1
eusilc$worker[eusilc$pl030 == "2"]<-1
eusilc$austrian<-0
eusilc$austrian[eusilc$pb220a == "AT"]<-1

# Computation, taking sample design into account
eusilc_many_prop <- many_prop(
  eusilc,
  list_vars = c(worker,austrian),
  list_vars_lab = c("% of workers", "% of Austrian"),
  facet = rb090,
  strata = db040,
  ids = db030,
  weight = rb050,
  title = "Proportion of workers and Autrian according to gender",
  subtitle = "Example with austrian SILC data from 'laeken' package"
)

# Results in graph form
```

```
eusilc_many_prop$graph
# Results in table format
eusilc_many_prop$tab
```

many_val_group	<i>many_val_group</i>
----------------	-----------------------

Description

Function to compare de proportions/means/medians of a set of several binary/quantitatives variables between different groups, based on complex survey data. It produces a list containing a table, including the confidence intervals of the indicators and a ready-to-be published ggplot graphic. The confidence intervals are taking into account the complex survey design.

Exporting the results to an Excell file is possible.

Usage

```
many_val_group(
  data,
  group,
  list_vars,
  type,
  list_vars_lab = NULL,
  facet = NULL,
  filter_exp = NULL,
  ...,
  na.rm.group = TRUE,
  na.rm.facet = TRUE,
  na.vars = "rm",
  total = TRUE,
  prop_method = "beta",
  position = "dodge",
  show_ci = TRUE,
  show_n = FALSE,
  show_value = TRUE,
  show_labs = TRUE,
  total_name = NULL,
  scale = NULL,
  digits = 0,
  unit = NULL,
  dec = NULL,
  pal = "OBSS_alt3",
  direction = 1,
  desaturate = 0,
  lighten = 0,
  darken = 0,
```

```

    dodge = 0.9,
    font = "Roboto",
    wrap_width_y = 25,
    wrap_width_leg = 25,
    legend_ncol = 4,
    title = NULL,
    subtitle = NULL,
    xlab = NULL,
    ylab = NULL,
    legend_lab = NULL,
    caption = NULL,
    lang = "fr",
    theme = "fonctionr",
    coef_font = 1,
    export_path = NULL,
    parallel = NULL
  )

many_prop_group(..., type = "prop")

many_median_group(..., type = "median")

many_mean_group(..., type = "mean")

```

Arguments

data	A dataframe or an object from the survey package or an object from the srvyr package.
group	A variable defining groups to be compared.
list_vars	A vector containing the names of the dummy/quantitative variables on which to compute the proportions/means/medians.
type	"prop" to compute proportions by group ; "mean" to compute means by group ; "median" to compute medians by group.
list_vars_lab	A vector containing the labels of the dummy/quantitative variables to be displayed on the graphic and in the table of result. Default uses the variable names in list_vars.
facet	A variable defining the faceting group.
filter_exp	An expression filtering the data, preserving the design. Notice that filter_exp works as <code>srvyr::filter()</code> : it excludes observations for which filter_exp results into NA. It is often the case when NA is present on one of the filter variables.
...	All options possible in <code>srvyr::as_survey_design()</code> .
na.rm.group	TRUE if you want to remove observations with NA on the group variable. FALSE if you want to create a group with the NA values for the group variable. Default is TRUE.

<code>na.rm.facet</code>	TRUE if you want to remove observations with NA on the facet variable. FALSE if you want to create a facet with the NA values for the facet variable. Default is TRUE.
<code>na.vars</code>	The treatment of NA values in variables (<code>list_vars</code>). "rm" removes NA separately in each individual variable, "rm.all" removes every individual that has at least one NA in one variable. Default is "rm".
<code>total</code>	TRUE if you want to compute a total, FALSE if you don't. Default is TRUE. Total is not displayed nor computed if <code>position = 'flip'</code> .
<code>prop_method</code>	Type of proportion method used to compute confidence intervals. See survey::svyciprop() for details. Default is beta method.
<code>position</code>	Position adjustment for the ggplot. Default is "dodge". Other possible values are "flip" and "stack". "dodge" means that groups are on the y axis and variables are in different colors, "flip" means that variables are on the y axis and groups are in different colors, and "stack" means that groups are on the y axis and variables are stacking with different colors. The latter is useful when the variables are component of a broader sum variable (e.g. different sources of income). If <code>position = 'flip'</code> , total is not displayed nor computed. If <code>position = "stack"</code> , confidence intervals are never shown on the graphic.
<code>show_ci</code>	TRUE if you want to show the error bars on the graphic. FALSE if you do not want to show the error bars. Default is TRUE. If <code>position = "stack"</code> , confidence intervals are never shown on the graphic.
<code>show_n</code>	TRUE if you want to show on the graphic the number of observations in the sample for each group and variable. The number can vary between variables if <code>na.vars = "rm"</code> . FALSE if you do not want to show this number. Default is FALSE.
<code>show_value</code>	TRUE if you want to show the proportions/means/median for each group and variable on the graphic. FALSE if you do not want to show the proportions/means/medians. Default is TRUE.
<code>show_labs</code>	TRUE if you want to show axes labels. FALSE if you do not want to show any labels on axes. Default is TRUE.
<code>total_name</code>	Name of the total bars on the graphic. Default is Total. Notice that total is not displayed nor computed if <code>position = 'flip'</code> .
<code>scale</code>	Denominator of the proportions. Default is 100 to interpret numbers as percentages. This argument is only used in case of <code>type = "prop"</code> .
<code>digits</code>	Number of decimal places displayed on the values labels on the graphic. Default is 0.
<code>unit</code>	Unit displayed on the graphic. Default is percent for <code>type = "prop"</code> and no unit for <code>type = "mean"</code> or <code>"median"</code> .
<code>dec</code>	Decimal mark displayed on the graphic. Default depends on lang: ", " for fr and nl ; "." for en.
<code>pal</code>	Colors of the bars. pal must be vector of R colors or hexadecimal colors or a palette from packages MetBrewer or PrettyCols or a palette from foncionr.
<code>direction</code>	Direction of the palette color. Default is 1. The opposite direction is -1.

desaturate	Numeric specifying the amount of desaturation where 1 corresponds to complete desaturation (no colors, grey layers only), 0 to no desaturation, and values in between to partial desaturation. Default is 0. See <code>colorspace::desaturate()</code> for details. If desaturate and lighten/darken arguments are used, lighten/darken is applied in a second time (i.e. on the color transformed by desaturate).
lighten	Numeric specifying the amount of lightening. Negative numbers cause darkening. Value should be ranged between -1 (black) and 1 (white). Default is 0. See <code>colorspace::lighten()</code> for details. If both argument lighten and darken are used (not advised), darken is applied in a second time (i.e. on the color transformed by lighten).
darken	Numeric specifying the amount of lightening. Negative numbers cause lightening. Value should be ranged between -1 (white) and 1 (black). Default is 0. See <code>colorspace::darken()</code> for details. If both argument lighten and darken are used (not advised), darken is applied in a second time (i.e. on the color transformed by lighten).
dodge	Width of the bars. Default is 0.9 to let a small space between bars. A value of 1 leads to no space between bars. Values higher than 1 are not advised because they cause an overlapping of the bars. <code>dodge</code> doesn't affect the spaces between sub-groups (group in case of <code>position = 'dodge'</code> or variables in case of <code>position = 'flip'</code>). There is always no space between sub-groups.
font	Font used in the graphic. See <code>load_and_active_fonts()</code> for available fonts. Default is "Roboto".
wrap_width_y	Number of characters before going to the line for the labels on de Y axe (groups if <code>position = 'dodge'</code> or <code>'stack'</code> , variables if <code>position = 'flip'</code>). Default is 25.
wrap_width_leg	Number of characters before going to the line for the labels the legend (variables if <code>position = 'dodge'</code> or <code>'stack'</code> , groups if <code>position = 'flip'</code>). Default is 25.
legend_ncol	Number maximum of columns in the legend. Default is 4.
title	Title of the graphic.
subtitle	Subtitle of the graphic.
xlab	X label on the graphic. As <code>ggplot2::coord_flip()</code> is used in the graphic, <code>xlab</code> refers to the x label on the graphic, after the <code>ggplot2::coord_flip()</code> , and not to the x variable in the data. Default (<code>xlab = NULL</code>) displays "Proportion : " (if <code>lang = "fr"</code>), "Proportion:" (if <code>lang = "en"</code>) or "Aandeel:" (if <code>lang = "nl"</code>) followed by the names of the variables (<code>list_vars</code>). To show no X label, use <code>xlab = ""</code> .
ylab	Y label on the graphic. As <code>ggplot2::coord_flip()</code> is used in the graphic, <code>ylab</code> refers to the y label on the graphic, after the <code>ggplot2::coord_flip()</code> , and not to the y variable in the data. Default (<code>ylab = NULL</code>) displays the name of the groups variable (if <code>position = 'dodge'</code> or <code>'stack'</code>) or no Y axe label (if <code>position = 'flip'</code>). To show no Y label, use <code>ylab = ""</code> .
legend_lab	Legend (fill) label on the graphic. Default (<code>legend_lab = NULL</code>) displays no legend label (if <code>position = 'dodge'</code> or <code>'stack'</code>) or the name of the groups variable (if <code>position = 'flip'</code>). To show no legend label, use <code>legend_lab = ""</code> .

caption	Caption of the graphic.
lang	Language of the indications on the graphic. Possibilities are "fr" (french), "nl" (dutch) and "en" (english). Default is "fr".
theme	Theme of the graphic. Default is "fonctionr". "IWEPS" adds y axis lines and ticks. NULL uses the default grey ggplot2 theme.
coef_font	A multiplier factor for font size of all fonts on the graphic. Default is 1. Usefull when exporting the graphic for a publication (e.g. in a Quarto document).
export_path	Path to export the results in an xlsx file. The file includes two sheets: the table and the graphic.
parallel	TRUE to enable parallel computing. Default is TRUE if replicated weights are used, otherwise default is FALSE.

Value

A list that contains a table and a graphic.

Examples

```
# Loading of data
data(eusilc, package = "laeken")

# Recoding variables
eusilc$worker <- 0
eusilc$worker[eusilc$pl030 == "1"] <- 1
eusilc$worker[eusilc$pl030 == "2"] <- 1
eusilc$austrian <- 0
eusilc$austrian[eusilc$pb220a == "AT"] <- 1

# Computation, taking sample design into account
eusilc_many_mean_group <- many_mean_group(
  eusilc,
  group = rb090,
  list_vars = c(py010n,py050n,py090n,py100n),
  list_vars_lab = c("Wage","Self-employment income","unemployment benefit","pension"),
  strata = db040,
  ids = db030,
  weight = rb050,
  title = "Average incomes according to gender",
  subtitle = "Example with austrian SILC data from 'laeken' package"
)

# Results in graph form
eusilc_many_mean_group$graph

# Results in table format
eusilc_many_mean_group$tab
```

official_pal	<i>official_pal</i>
--------------	---------------------

Description

A function that allows you to create the palettes of fonctionr.

Usage

```
official_pal(
  inst,
  n,
  direction = 1,
  desaturate = 0,
  lighten = 0,
  darken = 0,
  show_pal = FALSE,
  font = "Gotham Narrow",
  list_pal_names = FALSE
)
```

Arguments

<code>inst</code>	Name of the palette.
<code>n</code>	Number of colors.
<code>direction</code>	Direction of the palette color. Default is 1. The opposite direction is -1.
<code>desaturate</code>	Numeric specifying the amount of desaturation where 1 corresponds to complete desaturation, 0 to no desaturation, and values in between to partial desaturation.
<code>lighten</code>	Numeric specifying the amount of lightening. Negative numbers cause darkening.
<code>darken</code>	Numeric specifying the amount of lightening. Negative numbers cause lightening.
<code>show_pal</code>	TRUE to display a graph representing the specified color palette.
<code>font</code>	Font used in the graphic. See <code>load_and_active_fonts()</code> for available fonts.
<code>list_pal_names</code>	TRUE to generate a vector with palette names.

Value

A vector containing hexadecimal color codes.

Examples

```
official_pal("OBSS", 8, show_pal = TRUE)
official_pal("OBSS-Greens", 8, show_pal = TRUE)
official_pal("OBSS_div_mid4", 7, show_pal = TRUE)
official_pal("OBSS_div_bi3", 8, show_pal = TRUE)
official_pal("IBSA", 4, show_pal = TRUE)
official_pal("ULB", 6, show_pal = TRUE)
```

```
pivot_longer_survey  pivot_longer_survey
```

Description

Function to pivot, from wide to long, a dataframe produced by `srvyr::summarise()` with results by group(s).

Usage

```
pivot_longer_survey(data, n_groups)
```

Arguments

<code>data</code>	A dataframe produced by <code>srvyr::summarise()</code> with group(s).
<code>n_groups</code>	Number of groups by which data has been aggregated.

Value

A pivoted dataframe

Examples

```
# Loading data
data(eusilc, package = "laeken")

# Loading srvyr
library(srvyr)

# Making srvyr object
eusilc_srvyr <- as_survey_design(eusilc, ids = db030, strata = db040, weights = rb050)

# computing srvyr result using summarise()
result_srvyr <- eusilc_srvyr |>
  group_by(rb090, pb220a) |> # by sex and nationality
  summarise(mean_eqIncome = survey_mean(eqIncome), mean_age = survey_mean(age))

# Showing the srvyr summarise output
result_srvyr

# Pivoting the out with pivot_longer_survey()
```

```
pivoted_result <- pivot_longer_survey(result_srvyr, n_groups = 2)

# Output is pivoted
pivoted_result
```

prop_group	<i>prop_group</i>
------------	-------------------

Description

Function to compare a proportion among different groups based on complex survey data. It produces a list containing a table, including the confidence intervals of the indicators, a ready-to-be published ggplot graphic and a Chi-Square statistical test (using `survey::svychisq()`).

The confidence intervals and the statistical test are taking into account the complex survey design. In case of facets, the Chi-square test is computed on the total proportion between facets (and not within facets). In case of second group (`group.fill`), no Chi-square test is computed.

Exporting those results to an Excell file is possible.

Usage

```
prop_group(
  data,
  group,
  prop_exp,
  group.fill = NULL,
  facet = NULL,
  filter_exp = NULL,
  ...,
  na.rm.group = TRUE,
  na.rm.facet = TRUE,
  na.prop = "rm",
  total = TRUE,
  prop_method = "beta",
  reorder = FALSE,
  show_ci = TRUE,
  show_n = FALSE,
  show_value = TRUE,
  show_labs = TRUE,
  total_name = NULL,
  scale = 100,
  digits = 0,
  unit = "%",
  dec = NULL,
  col = "deepskyblue3",
  pal = "OBSS_Relax",
  direction = 1,
  desaturate = 0,
```

```

    lighten = 0,
    darken = 0,
    dodge = 0.9,
    font = "Roboto",
    wrap_width_y = 25,
    wrap_width_leg = 25,
    legend_ncol = 4,
    title = NULL,
    subtitle = NULL,
    xlab = NULL,
    ylab = NULL,
    legend_lab = NULL,
    caption = NULL,
    lang = "fr",
    theme = "fonctionr",
    coef_font = 1,
    export_path = NULL
  )

```

Arguments

data	A dataframe or an object from the survey package or an object from the srvyr package.
group	A variable defining the groups to be compared.
prop_exp	An expression defining the proportion to be computed. Notice that if <code>na.prop = "rm"</code> , <code>is.na()</code> is not allowed in this argument. The removal of NA is done before the computation of the proportion. Thus any function that takes into account NA (e.g. <code>%in%</code>) will not work as designed in this argument, unless <code>na.prop = "include"</code> .
group.fill	A variable defining a second variable of groups to be compared.
facet	A variable defining the faceting groups.
filter_exp	An expression filtering the data, preserving the design. Notice that <code>filter_exp</code> works as <code>srvyr::filter()</code> : it excludes observations for which <code>filter_exp</code> results into NA. It is often the case when NA is present on one of the filter variables.
...	All options possible in <code>srvyr::as_survey_design()</code> .
na.rm.group	TRUE if you want to remove observations with NA on the group and the <code>group.fill</code> variables. FALSE if you want to create a group with the NA values for the group variable and a <code>group.fill</code> with the NA values for the <code>group.fill</code> variable. Default is TRUE.
na.rm.facet	TRUE if you want to remove observations with NA on the facet variable. FALSE if you want to create a facet with the NA values for the facet variable. Default is TRUE.
na.prop	"rm" to remove observations with NA on one of the variables used in <code>prop_exp</code> before computing the proportions, "include" to compute the proportions with the NA in the denominators. Default is "rm". If <code>na.prop = "rm"</code> the function <code>is.na()</code> is not allowed in <code>prop_exp</code> .

total	TRUE if you want to compute a total, FALSE if you don't. The default is TRUE.
prop_method	Type of proportion method used to compute confidence intervals. See <code>survey::svyciprop()</code> for details. Default is beta method.
reorder	TRUE if you want to reorder the groups according to the proportion. NA value, if <code>na.rm.group = FALSE</code> , is not included in the reorder. In case of facets, the groups are reordered based on each median group. Default is FALSE.
show_ci	TRUE if you want to show the error bars on the graphic. FALSE if you don't want to show the error bars. Default is TRUE.
show_n	TRUE if you want to show on the graphic the number of observations in the sample in each group. FALSE if you don't want to show this number. Default is FALSE.
show_value	TRUE if you want to show the proportions in each group on the graphic. FALSE if you don't want to show the proportion. Default is TRUE.
show_labs	TRUE if you want to show axes and legend (in case of a <code>group.fill</code>) labels. FALSE if you don't want to show any labels on axes and legend. Default is TRUE.
total_name	Name of the total displayed on the graphic. Default is "Total" in French and in English and "Totaal" in Dutch.
scale	Denominator of the proportions. Default is 100 to interpret numbers as percentages.
digits	Number of decimal places displayed on the values labels on the graphic. Default is 0.
unit	Unit displayed on the graphic. Default is "%".
dec	Decimal mark displayed on the graphic. Default depends on lang: ", " for fr and nl ; "." for en.
col	Color of the bars if there is no <code>group.fill</code> . col must be a R color or an hexadecimal color code. Default is "deepskyblue3". The colors of total and NA group (in case of <code>na.rm.group = FALSE</code>) are always "grey40" and "grey". If there is a <code>group.fill</code> , col has no effect and pal argument should be used instead.
pal	Colors of the bars if there is a <code>group.fill</code> . pal must be vector of R colors or hexadecimal colors or a palette from packages MetBrewer or PrettyCols or a palette from fonctionr. The color of missing values in <code>group.fill</code> (in case of <code>na.rm.group = FALSE</code>) and of the total are always "grey" and "grey40". If there is no <code>group.fill</code> , pal has no effect and col argument should be used instead.
direction	Direction of the palette color. Default is 1. The opposite direction is -1. If there is no <code>group.fill</code> , this argument has no effect.
desaturate	Numeric specifying the amount of desaturation where 1 corresponds to complete desaturation (no colors, grey layers only), 0 to no desaturation, and values in between to partial desaturation. Default is 0. It affects only the palette (pal, if there is a second group) and not the monocolor (col, if there is no second group). See <code>colorspace::desaturate()</code> for details. If desaturate and lighten/darken arguments are used, lighten/darken is applied in a second time (i.e. on the color transformed by desaturate).

lighten	Numeric specifying the amount of lightening. Negative numbers cause darkening. Value should be ranged between -1 (black) and 1 (white). Default is 0. It doesn't affect the color of NA (in case of <code>na.rm.group = FALSE</code>). It affects only the palette (<code>pal</code> , if there is a second group) and not the monocolor (<code>col</code> , if there is no second group). See <code>colorspace::lighten()</code> for details. If both argument <code>lighten</code> and <code>darken</code> are used (not advised), <code>darken</code> is applied in a second time (i.e. on the color transformed by <code>lighten</code>).
darken	Numeric specifying the amount of lightening. Negative numbers cause lightening. Value should be ranged between -1 (white) and 1 (black). Default is 0. It doesn't affect the color of NA (in case of <code>na.rm.group = FALSE</code>). It affects only the palette (<code>pal</code> , if there is a second group) and not the monocolor (<code>col</code> , if there is no second group). See <code>colorspace::darken()</code> for details. If both argument <code>lighten</code> and <code>darken</code> are used (not advised), <code>darken</code> is applied in a second time (i.e. on the color transformed by <code>lighten</code>).
dodge	Width of the bars. Default is 0.9 to let a small space between bars. A value of 1 leads to no space between bars. Values higher than 1 are not advised because they cause an overlapping of the bars. <code>dodge</code> doesn't affect the spaces between second groups (<code>group.fill</code>). There is always no space between second groups.
font	Font used in the graphic. See <code>load_and_active_fonts()</code> for available fonts. Default is "Roboto".
wrap_width_y	Number of characters before going to the line for the labels of the groups. Default is 25.
wrap_width_leg	Number of characters before going to the line for the labels of the <code>group.fill</code> . Default is 25.
legend_ncol	Number of columns in the legend. Default is 4.
title	Title of the graphic.
subtitle	Subtitle of the graphic.
xlab	X label on the graphic. As <code>ggplot2::coord_flip()</code> is used in the graphic, <code>xlab</code> refers to the x label on the graphic, after the <code>ggplot2::coord_flip()</code> , and not to the x variable in the data. Default (<code>xlab = NULL</code>) displays "Proportion : " (if <code>lang = "fr"</code>), "Proportion:" (if <code>lang = "en"</code>) or "Aandeel:" (if <code>lang = "nl"</code>), followed by the <code>prop_exp</code> argument. To show no X label, use <code>xlab = ""</code> .
ylab	Y label on the graphic. As <code>ggplot2::coord_flip()</code> is used in the graphic, <code>ylab</code> refers to the y label on the graphic, after the <code>ggplot2::coord_flip()</code> , and not to the y variable in the data. Default (<code>ylab = NULL</code>) displays the name of the group variable. To show no Y label, use <code>ylab = ""</code> .
legend_lab	Legend (fill) label on the graphic. Default (<code>legend_lab = NULL</code>) displays the name of the <code>group.fill</code> variable. To show no legend label, use <code>legend_lab = ""</code> .
caption	Caption of the graphic. This caption goes under the default caption showing the result of the Chi-Square test. There is no way of not showing the result of the chi-square test as a caption.
lang	Language of the indications on the graphic. Possibilities are "fr" (french), "nl" (dutch) and "en" (english). Default is "fr".

theme	Theme of the graphic. Default is "fonctionr". "IWEPS" adds y axis lines and ticks. NULL uses the default grey ggplot2 theme.
coef_font	A multiplier factor for font size of all fonts on the graphic. Default is 1. Usefull when exporting the graphic for a publication (e.g. in a Quarto document).
export_path	Path to export the results in an xlsx file. The file includes three (without group.fill) or two sheets (with a group.fill): the table, the graphic and the Chi-Square statistical test result.

Value

A list that contains a table, a ggplot graphic and, in most cases, a Chi-square statistical test.

Examples

```
# Loading of data
data(eusilc, package = "laeken")

# Recoding eusilc$pl030 into eusilc$pl030_rec
eusilc$pl030_rec <- NA
eusilc$pl030_rec[eusilc$pl030 == "1"] <- "Working full time"
eusilc$pl030_rec[eusilc$pl030 == "2"] <- "Working part time"
eusilc$pl030_rec[eusilc$pl030 == "3"] <- "Unemployed"
eusilc$pl030_rec[eusilc$pl030 == "4"] <- "Student"
eusilc$pl030_rec[eusilc$pl030 == "5"] <- "Retired"
eusilc$pl030_rec[eusilc$pl030 == "6"] <- "Permanently disabled"
eusilc$pl030_rec[eusilc$pl030 == "7"] <- "Fulfilling domestic tasks"

# Computation, taking sample design into account
eusilc_prop <- prop_group(
  eusilc,
  group = pl030_rec,
  prop_exp = py090n > 0,
  strata = db040,
  ids = db030,
  weight = rb050,
  title = "% of ind. receiving unemployment benefits in their hh",
  subtitle = "Example with austrian SILC data from 'laeken' package"
)

# Results in graph form
eusilc_prop$graph

# Results in table format
eusilc_prop$tab
```

Description

Function to recode the default labels of a factor created by `cut()` from base R into more intuitive labels.

Usage

```
relab_cut(vec, suffix = NULL, right = TRUE, lang = "fr")
```

Arguments

<code>vec</code>	The vector to be recoded. It should be produced by <code>cut()</code> . Notice that the labels may not include scientific notation. Thus, in <code>cut()</code> , <code>dig.lab</code> argument should be high enough in order to produce labels without scientific notation.
<code>suffix</code>	The suffix to be indicated after the values. Usually, the unit of the variable will be used (e.g. euros, percents). Default is <code>NULL</code> , for no suffix.
<code>right</code>	<code>TRUE</code> if categories have been created with parameter <code>right = TRUE</code> in <code>cut()</code> . <code>FALSE</code> if categories have been created with parameter <code>right = FALSE</code> in <code>cut()</code> . Default is <code>TRUE</code> .
<code>lang</code>	Language of new labels. Possibilities are "fr" (french), "nl" (dutch) and "en" (english). Default is "fr".

Value

A vector with new labels.

Examples

```
cut(1:1000, breaks = 5, include.lowest = TRUE, right = FALSE, dig.lab = 4) |>
table()
```

```
cut(1:1000, breaks = 5, include.lowest = TRUE, right = FALSE, dig.lab = 4) |>
relab_cut(suffix = "€", right = FALSE) |>
table()
```

 theme_fonctionr

 theme_fonctionr

Description

A ggplot theme that is ready to use. It is used by most other functions, but can also be applied to an external ggplot object.

Usage

```
theme_fonctionr(  
  font = "Roboto",  
  theme = "fonctionr",  
  display = "ggplot",  
  grid.lines = "x",  
  coef_font = 1  
)
```

Arguments

font	Font used in the graphic. See <code>load_and_active_fonts()</code> for available fonts.
theme	The optionnal theme you want for the graphic. Available themes: "fonctionr" and "IWEPS". Default is NULL.
display	The way <code>theme_fonctionr()</code> works on the axis texts: like <code>ggplot2</code> or <code>ggtext</code> .
grid.lines	Specify major grid lines : "x", "y" or "both". Default is "x".
coef_font	A multiplier factor for font size.

Value

No return value, called for side effects.

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